```
logo
NTDB id 1124126 EL309 RS08465 WP 004283291.1  MKKILLVVAVSLVLGGCATDGTVDKDAQITQDWTVDRLYTEAQDELNSGNYTRAIKLYEILESRFPNGRYAQQSQLDTAY   80
NTDB id 1102 NMB RS03635 WP 002244053.1       MKKILLTVSLGLALSACATQGTVDKDAQITQDWSVEKLYAEAQDELNSSNYTRAVKLYEILESRFPTSRHAQQSQLDTAY   80
NTDB id 1106 NGFG RS01495 WP 003687645.1      MKKILLTVSLGLALSACATQGTADKDAQITQDWSVEKLYAEAQDELNSSNYTRAVKLYEILESRFPTSRHARQSQLDTAY   80
consensus                                     !!!!!!*!***!*!**!!!*!!*!!!!!!!!!!*!**!!*!!!!!!!!*!!!!!*!!!!!!!!!!!!**!*!*!!!!!!!!!

logo
NTDB id 1124126 EL309 RS08465 WP 004283291.1  AYYKDDEPEKALAAIDRFQRTHPRHPNLDYALYLKGLVLFNEDQSFLSKLASQDWSDRDPKANRDAYHAFDQLIKQYPNS  160
NTDB id 1102 NMB RS03635 WP 002244053.1       AYYKDDEKDKALAAIDRFRRLHPQHPNMDYALYLRGLVLFNEDQSFLNKLASQDWSDRDPKANREAYQAFAELVQRFPNS  160
NTDB id 1106 NGFG RS01495 WP 003687645.1      AYYKDDEKDKALAAIERFRRLHPQHPNMDYALYLRGLVLFNEDQSFLNKLASQDWSDRDPKANREAYQAFAELVQRFPNS  160
consensus                                     !!!!!!!**!!!!!!*!!*!*!!*!!!*!!!!!!*!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!*!!*!!**!****!!!

logo
NTDB id 1124126 EL309 RS08465 WP 004283291.1  KYVQDATERMAKLVDALGGNEIAVARYYMKRGAFLAAANRAQKVVQQYQNTRFVEEALAIMELAYIRLDNKQLAEDTHRI  240
NTDB id 1102 NMB RS03635 WP 002244053.1       KYAADATARMVKLVDALGGNEMSVARYYMKRGAYIAAANRAQKIIGSYQNTRYVEESLAILELAYKKLDKPRLAADTRRV  240
NTDB id 1106 NGFG RS01495 WP 003687645.1      KYAADATARMVKLVDALGGNEMSVARYYMKRGAYIAAANRAKKIIGSYQNTRYVEESLAILELAYKKLDKPQLAADTRRV  240
consensus                                     !!**!!!*!!*!!!!!!!!!!!**!!!!!!!!!!!**!!!!!!*!****!!!!!*!!!*!!!!*!!!!**!!***!!*!!*!*

logo
NTDB id 1124126 EL309 RS08465 WP 004283291.1  LEKNFPSSPYLTKPWQLDAMPWWRYWK   267
NTDB id 1102 NMB RS03635 WP 002244053.1       LETNFPKSPFLKQPWRSDDMPWWRYWH   267
NTDB id 1106 NGFG RS01495 WP 003687645.1      LETNFPKSPFLTHAWQPDDMPWWRYWH   267
consensus                                     !!*!!!*!!*!* *!* !*!!!!!!!*
```

X non conserved

X similar

X ≥ 50% conserved